

RESEARCH ARTICLE

The stage-specifically accelerated brain aging in never-treated first-episode patients with depression

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Abstract

Depression associated with structural brain abnormalities is hypothesized to be related with accelerated brain aging. However, there is far from a unified conclusion because of clinical variations such as medication status, cumulative illness burden. To explore whether brain age is accelerated in never-treated first-episode patients with depression and its association with clinical characteristics, we constructed a prediction model where gray matter volumes measured by voxel-based morphometry derived from T1-weighted MRI scans were treated as features. The prediction model was first validated using healthy controls (HCs) in two Chinese Han datasets (Dataset 1, $N = 130$ for HCs and $N = 195$ for patients with depression; Dataset 2, $N = 270$ for HCs) separately or jointly, then the trained prediction model using HCs ($N = 400$) was applied to never-treated first-episode patients with depression ($N = 195$). The brain-predicted age difference (brain-PAD) scores defined as the difference between predicted brain age and chronological age, were calculated for all participants and compared between patients with age-, gender-, educational level-matched HCs in Dataset 1. Overall, patients presented higher brain-PAD scores suggesting patients

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with depression having an “older” brain than expected. More specially, this difference occurred at illness onset (illness duration <3 months) and following 2 years then disappeared as the illness further advanced (>2 years) in patients. This phenomenon was verified by another data-driven method and significant correlation between brain-PAD scores and illness duration in patients. Our results reveal that accelerated brain aging occurs at illness onset and suggest it is a stage-dependent phenomenon in depression.

KEYWORDS

brain age, first-episode depression, machine learning, structural brain imaging

1 | INTRODUCTION

As a common mental disorder, depression that characterized by cognitive and affective deficits (Pan et al., 2018) leads to reduced quality of life and even suicide in patients with depression (S. Han et al., 2019). In addition to affective symptomatology, new evidences suggest that depressed individuals evidently present an increased risk of being attacked by aging-related somatic diseases (Nicholson, Kuper, & Hemingway, 2006) and mortality (Penninx, 2017) independent of suicide (Nock, Hwang, Sampson, & Kessler, 2010). These evidences include association between depression and decline in cognitive state (John, Patel, & Rusted, 2019), elevated risk of metabolic syndrome (Vancampfort et al., 2014) and cellular aging (Verhoeven et al., 2014) in patients with depression. Recent years, researchers began to use brain images combined with machine learning method to predict brain age (Gaser, Franke, Klöppel, Koutsouleris, & Sauer, 2013; Habes & Janowitz, 2016; Hajek et al., 2019; He et al., 2020) to explore disease such as schizophrenia, mild cognitive impairment (MCI). Exploring whether and how brain aging patterns are altered using machine learning could deepened our understanding of physiological mechanism of these disease.

As one of the brain imaging methods, magnetic resonance imaging (MRI) has the unique ability to noninvasively investigate brain structure and function. Previous structural MRI studies have identified pattern of neuroanatomical change along with normal brain development and aging at the individual level (Bashyam et al., 2020; Jylhävä, Pedersen, & Hägg, 2017). Recently, studies employ machine learning method combined with structural brain MRI images accurately predict individual brain age (Cole, Franke, & Cherbuin, 2019). Then, the prediction model is successfully applied to the study of several neurological diseases and reveals accelerated brain aging in disease such as schizophrenia, MCI, Alzheimer's disease, and autism (Gaser et al., 2013; Habes & Janowitz, 2016; Hajek et al., 2019; He et al., 2020). However, there is a paucity of studies exploring whether and how brain aging patterns is disturbed in patients with depression.

Although accelerated aging trajectories in patients with depression have been supported by mounts of evidences ranging from functional to biological state (Darrow et al., 2016; L. K. M. Han et al., 2018; Lever-van Milligen, Lamers, Smit, & Penninx, 2017;

Lindqvist et al., 2018), the number of studies specifically investing brain aging in depression with structural brain MRI is scarce. What is worse, the findings are not unified (L. K. M. Han, Dinga, Hahn, Ching, & Eyler, 2020). The reason might be the heterogeneity of samples clinical variations such as medicine, recurrence status (first vs. recurrent episode), age of onset, especially for illness duration (Besteher, Gaser, & Nenadić, 2019; L. K. M. Han et al., 2020; Kaufmann & van der Meer, 2019; Koutsouleris et al., 2014). In addition, three main questions remain unsolved in these studies. First, most of patients used in these studies are taking medicine (e.g., antidepressant) at the time of scan that changes brain structure (Cousins & Goodyer, 2015; Hamilton, Siemer, & Gotlib, 2008; Lavretsky, Roybal, Ballmaier, Toga, & Kumar, 2005; Willner, Scheel-Krüger, & Belzung, 2013), the effect of medicine cannot be completely eliminated. Second, patients enrolled in these studies are mainly Caucasian, whether the findings can be extended to other populations having different genetics, culture and environmental exposures remains unknown (S. Han & Ma, 2014; Nisbett & Miyamoto, 2005; Tang et al., 2018). The last and most important one, no study explores effect of cumulative illness burden on brain aging in depression. Gray matter (GM) volumes are found differently altered at different stages of the illness duration (Bora, Fornito, Pantelis, & Yücel, 2012; Serra-Blasco et al., 2013) that is thought to be one of the import potential confounders resulting inconsistent findings in depression (Chen et al., 2016). Prospective studies find that major depressive disorder with longer illness duration often experiences increasingly persistent disease with greater proportions of time spent in the midst of a major depressive episode (Judd et al., 1998; Solomon et al., 2000), suggesting neuropathological progression of depression. For example, right hippocampus is found reduced in older patients with depression particularly in patients with a longer course of illness (Bell-McGinty et al., 2002) and its volume decline is correlated with cumulative illness duration (Sheline, Gado, & Kraemer, 2003). The same progression of GM reduction is observed in rostral anterior cingulate cortex and dorsomedial frontal cortex (Frodl et al., 2008). Unfortunately, there is no study exploring the effect of illness duration on brain aging in depression.

To answer these questions, we constructed a prediction model where GM volumes were treated as features to explore whether brain age is accelerated and its association with clinical characteristics in

never-treated first-episode patients with depression. Gray matter volumes were quantified using voxel-based morphometry (VBM). The prediction model was built and validated using HCs in two Chinese Han datasets (Dataset 1, $N = 130$ for healthy controls [HCs] and $N = 195$ for patients with depression; Dataset 2, $N = 270$ for HCs) separately or jointly. Then the trained model was applied to explore brain aging in patients with depression. Brain-predicted age difference (brain-PAD) scores defined as the difference between predicted brain age and chronological age, were calculated in patients with depression and then compared with that in age-, gender-, educational level-matched HCs. A positive brain-PAD reflects accelerated maturation of brain while negative one represents delayed brain development in depression. Based on previous findings, we expected to find higher brain-PAD scores in patients than that in HCs and explored association between brain-PAD score and crucial clinical characteristics including illness duration, age of onset and gender.

2 | MATERIALS AND METHODS

2.1 | Sample

2.1.1 | Dataset 1

Patients with depression were recruited from outpatient services of Department of Psychiatry, The First Affiliated Hospital of Zhengzhou University, Zhengzhou, China from January 2015 to now. Patients were diagnosed according Diagnostic and Statistical Manual of Mental Disorders, Fourth Edition for depression, this procedure was done by one chief physician and one well-trained psychiatrist. The following inclusion criteria were employed: (a) the patients with depression must be first episode; (b) never taking any antidepressants (or antipsychotics) and any other antidepressant treatment such as psychological therapy and electric shock treatment; and (c) Han Chinese and right handedness. Patients with depression would be excluded if they met one of the following exclusion criteria: (a) comorbidity of other mental/psychotic disorders and (b) previous episodes of manic symptoms. The clinical states of the patients were evaluated using the 24/17-items Hamilton Depression (HAM-D) scale.

HCs were recruited from the community through poster advertisement. None of them presented a history of serious medical or neuropsychiatric illness or a family history of major psychiatric or neurological illness in their first-degree relatives. All HCs were Han Chinese and right handedness.

In addition, participants (both HCs and patients) included in the current study must meet the following exclusion criteria: (a) taking drugs such as anesthesia, sleeping, and analgesia in the past 1 month; (b) substance abuse; (c) a history of brain tumor, trauma, surgery, or other organic body disease; (d) suffering from cardiovascular diseases, diabetes, hypertension; (e) contraindications for MRI scanning including fixed dentures, metal braces, artificial heart valves, and other metal foreign bodies in the body; and (f) other structural brain abnormalities revealed by MRI scan.

Written informed consents were obtained from all participants before experiment. The study was approved by the research ethical committee of The First Affiliated Hospital of Zhengzhou University.

2.1.2 | Dataset 2

Another dataset come from Southwest University Adult Lifespan Dataset (SALD) study. This dataset was obtained from healthy participants ($N = 494$, 308 female, 187 male, age range 19–80). The exclusion criteria included MRI-related exclusion criteria, current psychiatric/neurological disorders, use of psychiatric drugs in the past 3 months prior to scanning and so on. More detailed description about the subject information and data acquisition parameters, please see (Wei et al., 2018), 3D structural MRI sequence was in supplement results. Participants under the age of 50 ($N = 270$) were included in this study. The data is available for research purposes through the International Data-sharing Initiative (https://fcon_1000.projects.nitrc.org/indi/retro/sald.html).

2.2 | VBM analysis

All scans were processed using the CAT12 toolbox (<https://dbm.neuro.uni-jena.de/cat12/>). The standard pipeline steps were used including bias-field correction, segmentation (GM and white matter and cerebrospinal fluid, adjustment for partial volume effects, normalization into Montreal Neurological Institute space, resampled to $1.5 \text{ mm} \times 1.5 \text{ mm} \times 1.5 \text{ mm}$ and nonlinear modulation (Ashburner, 2009). Finally, the GM maps were smoothed using 6 mm full width at half maximum Gaussian kernel. The total intracranial volume (TIV) of each participant was also calculated for the next comparative analysis.

2.3 | Prediction model

The GM volumes of the whole brain were segmented into different regions using brain atlas dividing the whole brain into 268 regions in virtue of a group-wise spectral clustering algorithm (Finn et al., 2015; Shen, Tokoglu, Papademetris, & Constable, 2013). The mean GM volume of each region was used as features and sent into prediction model. Gaussian process regression (GPR) was used to predict participants' age from the mean GM volumes of all brain regions (Marquand, Rezek, Buitelaar, & Beckmann, 2016; Seeger, 2004). This approach was highly flexible and could accommodate nonlinear relationships delivering state-of-the-art prediction performance in many types of neuroimage data (Hyun et al., 2014). The GPR method was implemented in the Gaussian processes for machine learning (GPML) toolbox (www.gaussianprocess.org/gpml/code/). The model parameters were optimized using a conjugate gradient optimizer (also included in GPML toolbox) as done in before (Marquand et al., 2016; Marquand et al., 2016). We also compared the performance of different machine learning methods commonly used in previous studies (He et al., 2020;

Koutsouleris et al., 2014; Sone et al., 2019). Specially, we compared the prediction performance of different machine learners (GPR vs. support vector regression [SVR]) and feature selection strategies (mean GM volumes based on atlas vs. principal component analysis [PCA] (Franke, Ziegler, Klöppel, & Gaser, 2010)). When PCA was used, the first N eigenvariates explaining the 95% of variance were extracted as features in the training process.

2.4 | Model validation

Ten folds cross-validation (CV) were used to evaluate the performance of the prediction model (Sone et al., 2019; ZIEGEL, 2010). This procedure divided the samples into 10 subsets of (approximately) equal size. We trained the model 10 times where one of the subsets was treated as test sample set and the others train sample each time. The trained model learned from the train sample set was then applied to the test sample to obtain brain age. We calculated (a) mean absolute error (MAE) between estimated brain age (output of the prediction model) and chronological age and (b) the correlation between the chronological age and estimated brain age through 10-fold CV.

In this procedure, two datasets were used separately or jointly to evaluate the performance and universality of our prediction model. That is to say, 10-fold CV was done in dataset 1 (HCs, $N = 130$), dataset 2 ($N = 270$) and a collection of both datasets (HCs in Dataset 1 and all participants in dataset 2, $N = 400$) to evaluate the performance of the prediction model.

To eliminate the effect of chronological age on the statistical results of altered brain-PAD scores in patients as much as possible, we only included participants under age of 36 ($N = 182$) in Dataset 2 in the final prediction model built to compare brain-PAD scores between HCs and patients because that the brain-PAD scores were found to be highly related with chronological age (Hajek et al., 2019; L. K. M. Han et al., 2018; L. K. M. Han et al., 2020). Based on the prediction model, we calculated each participant's brain-PAD (score: predicted age–chronological age).

2.5 | Statistics

The brain-PAD scores were compared between patients with depression and HCs in Dataset 1 using two-tailed two-sample t test. Gender, age, and age² were used as covariates in statistical models (L. K. M. Han et al., 2020). Moreover, we also divided patients with depression into three stages according to illness duration (Stage 1: 0–12 months; Stage 2: 12–24 months; Stage 3: $> = 24$ months) to explore whether aberrance of brain-PAD scores was stage dependent, because mental disorders including depression were found to be related with progressive brain structural alterations (Cao, Passos, Mwangi, Amaral-Silva, & Tannous, 2017; Koutsouleris et al., 2014; Treadway et al., 2015; Yüksel et al., 2018; Zhang et al., 2017). At the same time, to explore whether the altered brain-PAD scores occurred at illness onset, we also compared brain-PAD scores in patients with much shorted illness

duration ($< 3/6$ months) with that in matched HCs. All above-mentioned statistical procedures were done between patients with depression and age-, gender-, and educational level-matched HCs in Dataset 1 and statistical results were adjusted for gender, age, and age².

Meanwhile, a newly proposed data-driven algorithm (HYDRA) (Chand et al., 2020; Varol, Sotiras, & Davatzikos, 2017) was used to divide patients with depression into subgroups according to structural GM volumes that used in the prediction model (details in supplement results). This procedure was done to validate stage-specific aberrance of brain-PAD scores in patients from the opposite direction. That is to say, if brain-PAD scores were stage-specifically altered, patients would be divided into subgroups showing significant difference of illness duration. HYDRA could perform classification and subtyping simultaneously where classification was performed through the separation of HCs from patients with depression and subtyping was carried out by clustering patients with hyperplanes. The number of subgroups was set 1 to 5. Finally, we explored difference of clinical variables including gender, age, age of first onset, and illness duration among subgroups.

Within patients with depression, we also consider the effect of factors such as gender (male vs. female), age of onset of depression (categorized as: early/adolescent onset, $< = 25$ years (Penttilä et al., 2009; Truong et al., 2013) (or 21 years (Schmaal et al., 2016; Schmaal et al., 2017)) years; middle adulthood/adult onset, > 25 years (or 21) years) on the brain-PAD scores. Especially, to explore the difference of brain-PAD scores between male and female patients, we built prediction model using male and female HCs jointly or separately. In this procedure, patients were divided into two subgroups according to factors (male vs. female or adolescent onset vs. adult onset). Then, brain-PAD scores were compared between subgroups using two sample t test adjusted for gender (not for gender difference), age and age². We also compared the TIV of patients with that of HCs at each stage and across stages.

To explore the relationship between brain-PAD scores and clinical variables, we calculated Spearman correlations between brain-PAD scores and the total HAMD score and illness duration.

3 | RESULTS

3.1 | Clinical demographics

The clinical demographics are presented in Table 1. The distribution of participants' age is drawn in Figure 1. The clinical demographics of patients at different stages and matched HCs were in Table S1.

3.2 | Prediction model performance in HCs

Within healthy participants, our proposed prediction model presented steady performance across datasets (Figure 1). We also found performance of the prediction model was better when the sample size was

TABLE 1 Demographic and clinical characteristics of participants

	Dataset 1		Dataset 2
	HC (N = 130)	Depression (N = 195)	Subjects (N = 270)
Male, No. (%)	59 (45.38)	95 (48.7)	98 (36.30)
Age, mean (SD) [range], years	21.25 (5.33) [12–36]	18.14 (4.47) [11–37]	31.50 (9.99) [19–50]
Educational level, mean (SD), years	13.56 (4.50)	10.11 (2.13)	—
Duration of illness, mean (SD), months	—	15.74 (16.96)	—
HAMD score, mean (SD), [range]	—	22.38 (5.72) [12–48] ^a 39.29 (11.68) [20–61] ^b	—
Handedness, right/left	130/0	195/0	—
Age of first onset, years	—	16.81 (4.40)	—

Abbreviations: HAMD, Hamilton rating scale for depression, HC, healthy controls.

^a17-items HAMD for 167 patients.

^b24-items HAMD for 28 patients.

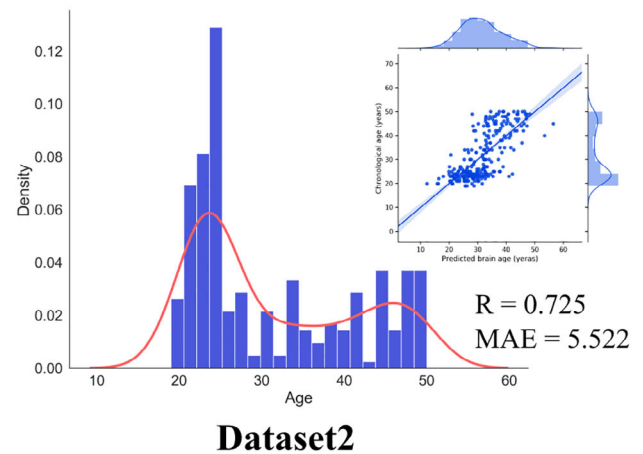
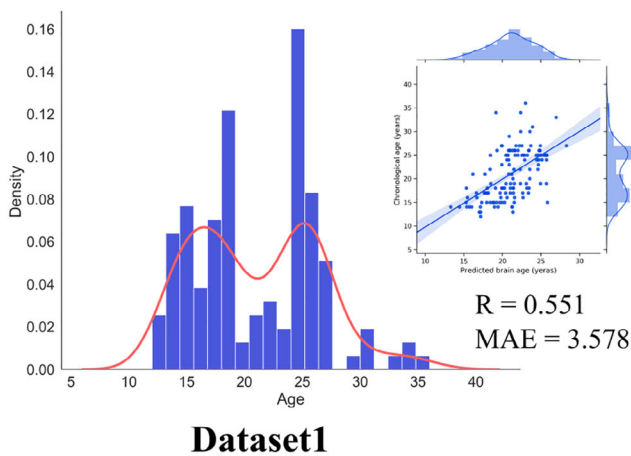
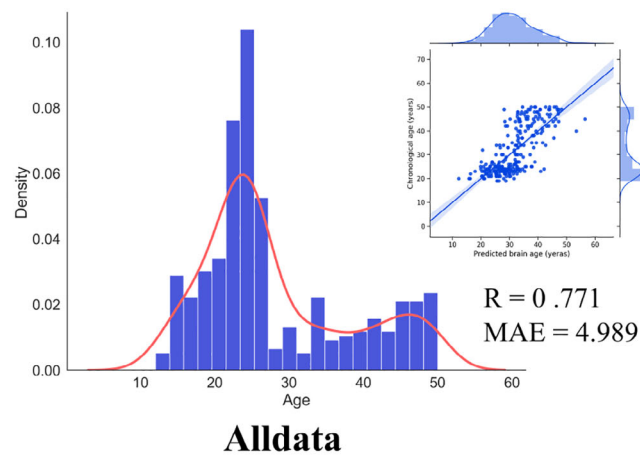


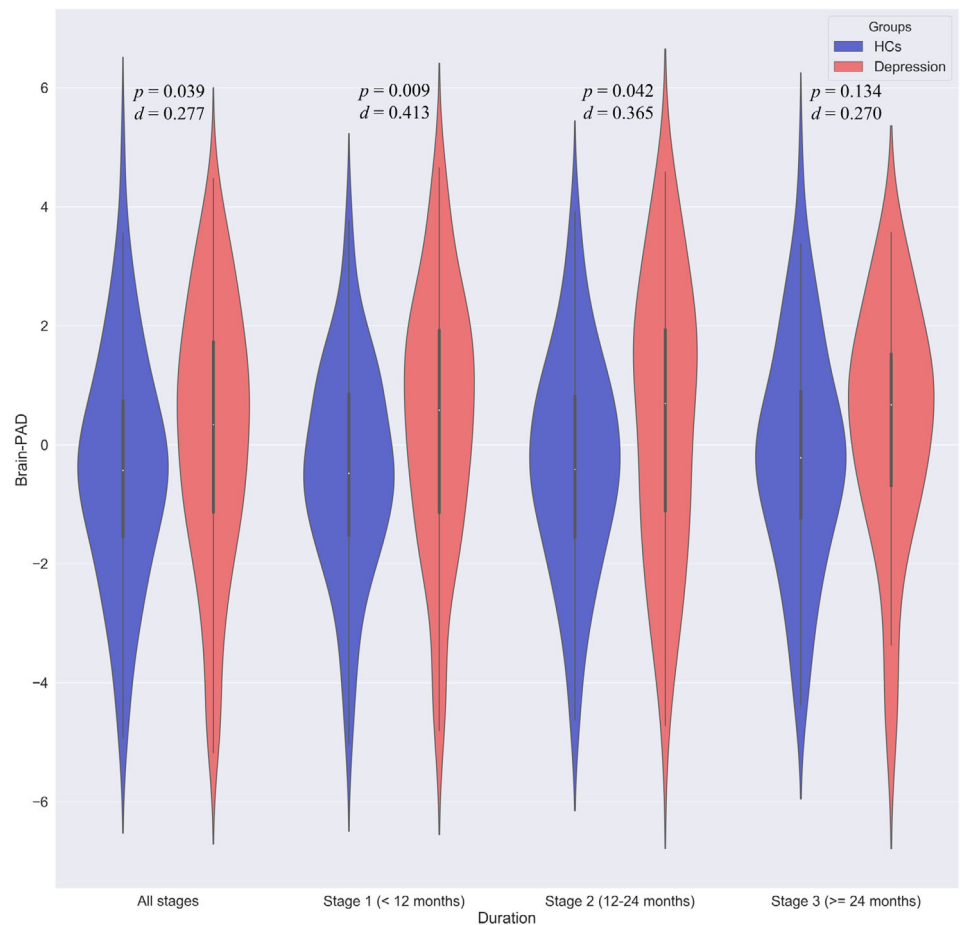
FIGURE 1 The distribution of age in datasets and the performance of the proposed prediction model. *R*: Pearson's correlation; MAE, mean absolute error

larger. This result was in line with the conclusion of previous study (Franke et al., 2010). In addition, we compared our proposed prediction model with other models commonly used in previous studies. As a result, the prediction model used in the current study presented overall better performance than other methods (Table S2).

3.3 | Brain-PAD scores in patients with depression

Overall, patients with depression presented higher brain-PAD scores than that in matched HCs (+0.586 years, $t = 2.065$, $p = 0.039$, Cohen's $d = 0.277$) adjusted for gender, age, and age².

FIGURE 2 Aberrance of brain-predicted age difference (brain-PAD) scores in patients at different stages adjusted for gender, age, and age²



Then, we divided patients into three stages according to illness duration and compared their brain-PAD scores with that in matched HCs. We found that only at the first two stage (Stage 1, +0.812 years, $t = 2.639$, $p = .009$, $d = 0.413$; Stage 2, +0.742 years, $t = 2.056$, $p = .042$, $d = 0.365$) presented significant higher brain-PAD scores than HCs, while patients with depression at Dstage 3 (illness duration $> = 24$ months) did not ($t = 1.507$, $p = .134$, $d = 0.270$). We further explored whether brain-PAD scores were higher in depression at illness onset. Brain-PAD scores of patients whose illness duration was $< 3/6$ months were compared with that in matched HCs. As a result, we found that patients with depression presented higher brain-PAD scores than that in HCs at the beginning of illness (< 3 months, +1.061 years, $t = 2.520$, $p = .013$, $d = 0.546$; < 6 months, +0.942 years, $t = 2.829$, $p = .005$, $d = 0.505$) (Figure 2).

As a complement to these results, we employed a data-driven method dividing patients into subgroups and then compared clinical variables among subgroups. The largest adjusted rand index (ARI) was found at $K = 2$ (ARI = 0.885) (Figure S1 in supplement results), suggesting patients with depression could be divided into two distinct subgroups. Then, we compared clinical variables including gender, age, age of first onset and illness duration between two subgroups. As a result, only illness duration presented significant difference between two subgroups ($t = -2.101$, $p = .0367$, $d = -0.301$) (Figure S2 in supplement results) (Figure 3).

There was no not significant difference of brain-PAD scores between female and male patients with depression, all p values $> .05$ in two different prediction models constructed using female and male participants separately or jointly. Performance of the prediction model using female and male participants separately was drawn in Figure S3 (Supplement results). We also did not find significant effect of age of onset on the brain-PAD scores in patients with depression.

Meanwhile, brain-PAD scores were significant correlated with illness duration in patients with depression (Spearman $R = -1.145$, $p = .043$). There was no significant correlation between brain-PAD scores with the total score of HAMD. We did not observe significant difference of TIV between patients and HCs at each stage and across stages (Figure 4).

4 | DISCUSSION

Patients with depression presented higher brain-PAD scores than HCs suggesting patients with depression having an “older” brain than expected. This phenomenon occurred since the beginning of the illness ($< 3/6$ months) and lasted for up to the first 2 years of illness. Meanwhile, results of HYDRA revealed two distinct subgroups of patients presenting characterized with different length of illness duration. Brain-PAD scores were negatively correlated with illness

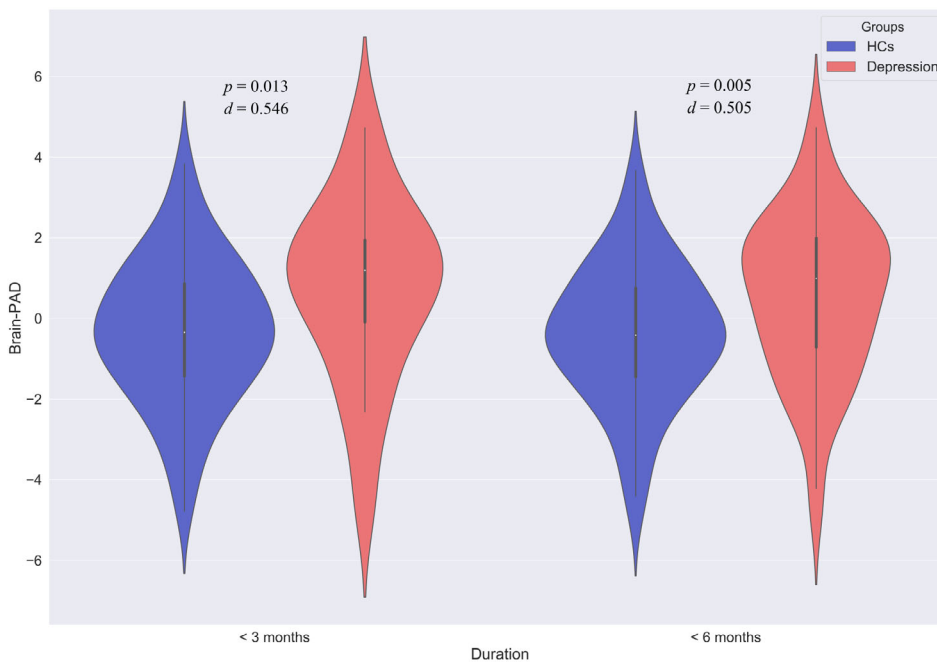


FIGURE 3 Aberrance of brain-predicted age difference (brain-PAD) scores in patients at illness onset (<3/6 months) adjusted for gender, age, and age²

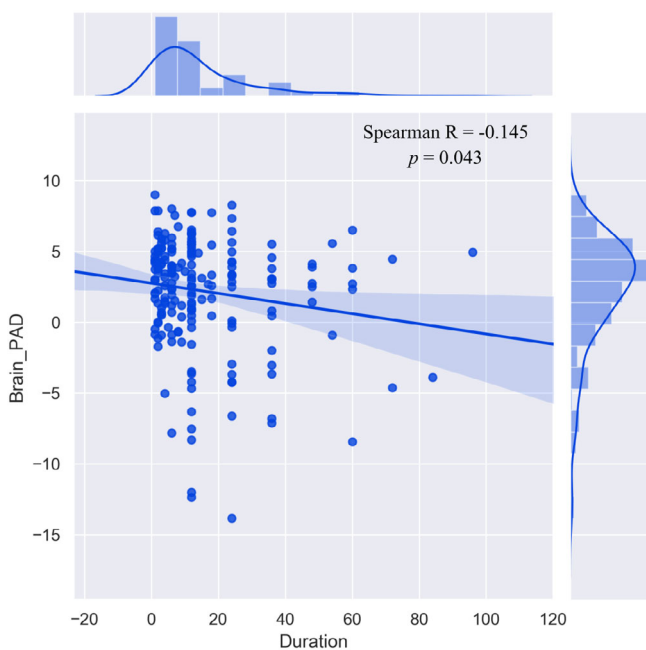


FIGURE 4 The correlation between brain-predicted age difference (brain-PAD) scores and illness duration in patients with depression

duration in patients with depression. These converging results confirmed accelerated brain aging occurred at illness onset and hinted it was a stage-dependent phenomenon in depression.

Our proposed prediction model presented stable performance across highly heterogeneous datasets having different scanning parameters and age distribution. Consistent with the conclusion of previous study focusing on influence of various parameters on performance of prediction model (Franke et al., 2010), we also found that

the performance of prediction model got better as the sample size increased in the results. The MAE (4.990 year in age range of 11–50 years, 0.128 after scaled to covered age range) of our proposed model built using all HCs across two datasets ($N = 400$) was close to that reported in previous studies (L. K. M. Han et al., 2020; Schnack et al., 2016). The comparatively large MAE in our results might be ascribed more to relatively small sample size. In addition, we compared different machine learners (GPR vs. SVR) and feature selection strategies (mean GM volumes based on atlas vs. PCA), the results illustrated that our proposed model had obvious better performance (Table S1 in supplement results) suggesting GPR was highly flexible delivering state-of-the-art prediction performance in many types of neuroimage data (Hyun et al., 2014). A more elaborate methodological comparison was omitted because it was beyond the scope of the current study.

As we hypothesized, brain-PAD scores were higher (+0.586 years for all, ranging from +0.742 to +1.061 years for different stages) in never-treated first-episode patients with depression than that in HCs. These results replicated earlier results (L. K. M. Han et al., 2020; Koutsouleris et al., 2014) in Chinese Han patients with depression suggesting the higher brain-PAD score in depression was not about race having different genetics, culture and environmental exposures. Higher brain-PAD scores suggesting accelerated aging trajectories were related to greater cognitive impairment in depression (Hatton et al., 2018). One proposed explanation was the common biological mechanism underlying depression and brain aging (Franceschi et al., 2000; L. K. M. Han et al., 2020), supported by findings that brain-PAD was temporarily reduced by 1.1 years introduced by the probable acute anti-inflammatory effects of ibuprofen in HCs (Le et al., 2018). Moreover, brain-PAD scores were higher even at illness onset (with illness duration <3 months) and not associated with clinical variations including gender and onset of illness in depression,

hinting it was related to pathomechanism of depression not the consequence of illness progression. Brain-PAD difference in our results was closer to that (+1.1 years) reported in Han et al. (L. K. M. Han et al., 2020), the effect size of difference was much higher than that reported before (L. K. M. Han et al., 2020), the reason might be that we only enrolled never-treated first-episode patients in the current study excluding effect of clinical variations such as medication status and acquisition protocols. Although the relatively smaller brain-PAD scores might partly come from factors including age range and distribution, scanning parameters and sample size, the more likely reason might be that higher brain age was a stage-dependent phenomenon in depression.

The difference of brain-PAD scores were initially prominent and then become less pronounced after 2 years of illness in patients, hinting higher brain age was a stage-dependent phenomenon in depression. Mounts of evidences suggested depression was a neuroprogressive illness (Moylan, Maes, Wray, & Berk, 2013). Longer and more frequent depressive episodes were accompanied by increase vulnerability to further relapses and functional decline (Kendler, Thornton, & Gardner, 2001). In a recent review (Moylan et al., 2013), the authors collected evidences covering from clinical, biochemical and neuroimaging studies and indicated neuroprogressive process occurring in depression. For example, GM volumes of the hippocampus and medial frontal cortex were associated with greater number of prior depressive episodes (Belleau, Treadway, & Pizzagalli, 2019), resulted from chronic stress and toxic effects of recurrent depressive episodes (Treadway et al., 2015) and the same reason leading to progressive anterior cingulate cortex and dorsomedial frontal changes in depression (Bora et al., 2012). In the latest research of Elaine et al., the authors found strong correlation between increasing duration of untreated illness and greater microglial activation, providing a new evidence of neuroprogression (Setiawan et al., 2018). Consistent with these studies, we found the differences of brain-PAD scores between patients with depression and HCs were stage-dependent. Specially, the difference of brain-PAD scores between patients and HCs were initially high and then become less pronounced after 2 years of illness in patients with depression. This phenomenon could mirror the transition from a clinically unstable period, with large variability in functioning, to a relatively stable period, when patients have reached a plateau in functioning (Davidson & McGlashan, 1997; van Haren et al., 2003). Notably, insignificant brain-PAD scores in patients with longer illness duration ($> = 2$ years) did not necessarily declare the remission of severity of illness, we did observe significant difference of the total score of HAMD ($p = .185$, $t = 1.331$) between patients with longer duration ($> = 2$ years) and ones with shorter illness duration (< 2 years). One reason of Bianca et al. failing to find accelerated aging in patients with depression might be illness duration of patients recruited was too long (Besteher et al., 2019). Although the illness duration of patients was not reported, the most of patients used in their study were experiencing multiple episodes tending to have a longer course of illness. Unfortunately, there were rare studies having complete information on the cumulative illness duration across episodes that was different from the index episode being a recurrent or first-

onset episode (Ten Have et al., 2017), given the relapsing and remitting nature of depression (Schmaal et al., 2016). More longitudinal studies with first-episode patients (or accurately recording cumulative disease burden) were needed to validate this assumption. In addition, we also found that patients were divided into two subgroups only having significantly different length of illness duration with data-driven algorithm (HYDRA). This result was consistent with another study also finding illness duration was an important factor varied between two subgroups revealed by cluster analysis (Corponi & Anmella, 2020). Previous studies also found patients with longer illness duration were accompanied by greater severity (Spijker et al., 2002; Ten Have et al., 2017), higher comorbidities' burden, suicide behavior (Spijker et al., 2002) and lower probability of recovery (Corponi & Anmella, 2020; Keller et al., 1992). Taken together, these findings convergently hinted accelerated brain aging was also related to illness duration and a stage-dependent phenomenon in depression.

There were several limitations must be considered. First, patients all come from a single dataset, another independent dataset was needed to validate our results in the future. Second, patient enrolled in the current study were under depressive state, whether the accelerated brain aging was trait related or state related should be tested with patients under the remitted state in future study (Rive et al., 2015; Van Eijndhoven et al., 2013). Third, the age of patients was less than 36 years, whether the conclusion held true for late-onset depression needed to be explored (Penttilä et al., 2009; Truong et al., 2013).

5 | CONCLUSION

In the current study, we enrolled Chinese Han patients to confirm accelerated brain aging and explore effects of clinical variations on the brain age in depression. Overall, patients presented higher brain-PAD scores suggesting patients with depression having an "older" brain than expected. This difference occurred in patients at illness onset (illness duration < 3 months) and following 2 years then disappeared as the illness further advances (> 2 years). In addition, we found that patients were divided into two subgroups characterized with length of illness duration, by data-driven method employed to explore factors affecting brain-PAD scores and validate our results. Combing with the correlation between brain-PAD scores and illness duration in patients, these converging results confirmed higher brain age and hinted accelerated brain aging was a stage-dependent phenomenon in depression.

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CONFLICT OF INTERESTS

The authors declare no conflict of interest.

DATA AVAILABILITY STATEMENT

Data sharing is not applicable to this article as no new data were created or analyzed in this study.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of this article.

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